

5'	9	18	27	36	45	54
GCC	GAA	CCC	CGC	CCA	CTC	GCT
GAA	CCC	CGC	CCA	CTC	GCT	GAA
AGT	GAA	GGA	GGA	GGA	GAA	AGT
GGC	GAG	GGA	GGA	GGA	GGA	GAG
	63	72	81	90	99	108
TTC	ATC	CCT	GCC	TAG	CGC	CCA
TTC	ATC	CCT	GCC	TAG	CGC	CCA
ATC	CCT	GCC	TAG	CGC	CCA	ATC
ATC	CCT	GCC	TAG	CGC	CCA	ATC
	117	126	135	144	153	162
GAT	GTG	GCG	ACT	GCG	GCC	GAC
GAT	GTG	GCG	ACT	GCG	GCC	GAC
CTG	GAT	GTG	GCG	ACT	GCG	GCC
CTG	GAT	GTG	GCG	ACT	GCG	GCC
	171	180	189	198	207	216
CAT	GAG	ATT	TTC	ACG	TCG	CCG
CAT	GAG	ATT	TTC	ACG	TCG	CCG
CTG	GAT	GTG	GCG	ACT	GCG	GCC
CTG	GAT	GTG	GCG	ACT	GCG	GCC
	225	234	243	252	261	270
CTC	TAC	AAG	ATC	GTG	CGC	GGG
CTC	TAC	AAG	ATC	GTG	CGC	GGG
CTC	TAC	AAG	ATC	GTG	CGC	GGG
CTC	TAC	AAG	ATC	GTG	CGC	GGG
	279	288	297	306	315	324
ACG	ACG	ANG	CCG	CCC	CCT	CTG
ACG	ACG	ANG	CCG	CCC	CCT	CTG
ACG	ACG	ANG	CCG	CCC	CCT	CTG
ACG	ACG	ANG	CCG	CCC	CCT	CTG
	333	342	351	360	369	378
GAG	CTG	CGG	CGC	TTC	GAC	GGC
GAG	CTG	CGG	CGC	TTC	GAC	GGC
GAG	CTG	CGG	CGC	TTC	GAC	GGC
GAG	CTG	CGG	CGC	TTC	GAC	GGC

FIG. 1A

387	396	405	414	423	432
AAC GGC AAG GTG TTT GAT GTG ACC AAA GGC CGC AAA TTC TAC GGG CCC GAG GGG					
N G K V F D V T K G R K F Y G P E G					
441	450	459	468	477	486
CCG TAT GGG GTC TTT GCT GGA AGA GAT GCA TCC AGG GGC CTT GCC ACA TTT TGC					
P Y G V F A G R D A S R G L A T F C					
495	504	513	522	531	540
CTG GAT AAG GAA GCA CTG AAG GAT GAG TAC GAT GAC CTT TCT GAC CTC ACT GCT					
L D K E A L K D E Y D D L S D L T A					
549	558	567	576	585	594
GCC CAG CAG GAG ACT CTG AGT GAC TGG GAG TCT CAG TTC ACT TTT AAG TAT CAT					
A Q Q E T L S D W E S Q F T F K Y H					
603	612	621	630	639	648
CAC GTG GGC AAA CTG CTG AAG GAG GAG GAG GGG GAG CCC ACT GTG TAC TCA GAT GAG					
H V G K L L K E G G E E P T V Y S D E					
657	666	675	684	693	702
GAA GAA CCA AAA GAT GAG AGT TCC CGG AAA AAT GTT AAA GCA TTC AGT GGA AGT					
F E P K D E S S S R K N V K A F S G S					
711	720	729	738	747	756
ATA TCT ATN NTG TAT TTT GCA AAA TCA TTT GTA ACA GTC CAC TNT GTC TTT AAA					
I S X X Y F A K S F V T V H X V F K					

FIG. 1B

765 ACA TAG TGT TAC AAT AAT TAG AAA GTT TGA GC 3'
774
783
'T

FIG. 1C

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Inventors: Goli et al
Title: NOVEL HUMAN CYTOKINE/STEROID RECEPTOR PROTEIN
Serial No.: To Be Assigned

1	M A A E D V V A T G A D P S <u>D</u> L E <u>S</u> G G	2504333
1	M A A E D V <u>V</u> A T G A D P S <u>E</u> L E G G G	GI 1518818
1	M A A E D V <u>A</u> <u>A</u> T G A D P S <u>E</u> <u>L</u> <u>E</u> G <u>G</u> G	GI 1657409
21	L L <u>H</u> E I F T S P L N L L L L G L C I F	2504333
21	L L <u>Q</u> E I F T S P L N L L L L G L C I F	GI 1518818
21	L L H E I F T S P L N L L L L G L C I F	GI 1657409
41	L L Y K I V R G D Q P <u>A</u> A S G D <u>R</u> T T T	2504333
41	L L Y K I V R G D Q P <u>G</u> A S <u>G</u> D N D D D	GI 1518818
41	L L Y K I V R G D Q P A A S - <u>D</u> S D D D	GI 1657409
61	X P P P L P R L K <u>R</u> R D F T P A E L R R	2504333
61	E P P P L P R L K <u>P</u> R D F T P A E L R R	GI 1518818
60	E P P P L P R L K R R D F T P A E L R R	GI 1657409
81	<u>F</u> D G V Q D P R I L M A I N G K V F D V	2504333
81	<u>Y</u> D G V Q D P R I L M A I N G K V F D V	GI 1518818
80	<u>F</u> D G V Q D P R I L M A I N G K V F D V	GI 1657409
101	T K G R K F Y G P E G P Y G V F A G R D	2504333
101	T K G R K F Y G P E G P Y G V F A G R D	GI 1518818
100	T K G R K F Y G P E G P Y G V F A G R D	GI 1657409
121	A S R G L A T F C L D K E A L K D E Y D	2504333
121	A S R G L A T F C L D K E A L K D E Y D	GI 1518818
120	A S R G L A T F C L D K E A L K D E Y D	GI 1657409
141	D L S D L T <u>A</u> A Q Q E T L <u>S</u> D W <u>E</u> S Q F	2504333
141	D L S D L T <u>P</u> A Q Q E T L <u>N</u> D W <u>D</u> S Q F	GI 1518818
140	D L S D L T <u>P</u> A Q Q E T L <u>N</u> D W <u>D</u> S Q F	GI 1657409
161	<u>T</u> F K Y H H V - - G K L L K E G E E P T	2504333
161	<u>S</u> S P S S T I T W G K L L <u>E</u> G A E E P I	GI 1518818
160	<u>T</u> F K Y H H V - - G K L L K E G E E P T	GI 1657409

FIG. 2A

179	V Y S D E E E P K D E S S R K N V K A F	2504333
181	V Y S D <u>D</u> E E <u>Q</u> K <u>M R L L G R V T E</u> <u>A</u> V	GI 1518818
178	V Y S D E E E P K D E S <u>A</u> <u>R</u> - - - - -	GI 1657409
199	<u>S G S I S</u> X X <u>Y F A K S F V T V H</u> X <u>V F</u>	2504333
201	<u>S G</u> <u>A Y L F L</u> <u>Y F A K S F V T</u> <u>F Q S</u> <u>V F</u>	GI 1518818
192	- - - - - K N - - - - - D	GI 1657409
219	<u>K T</u>	2504333
221	<u>T</u> <u>T</u> W	GI 1518818
194		GI 1657409

FIG. 2B

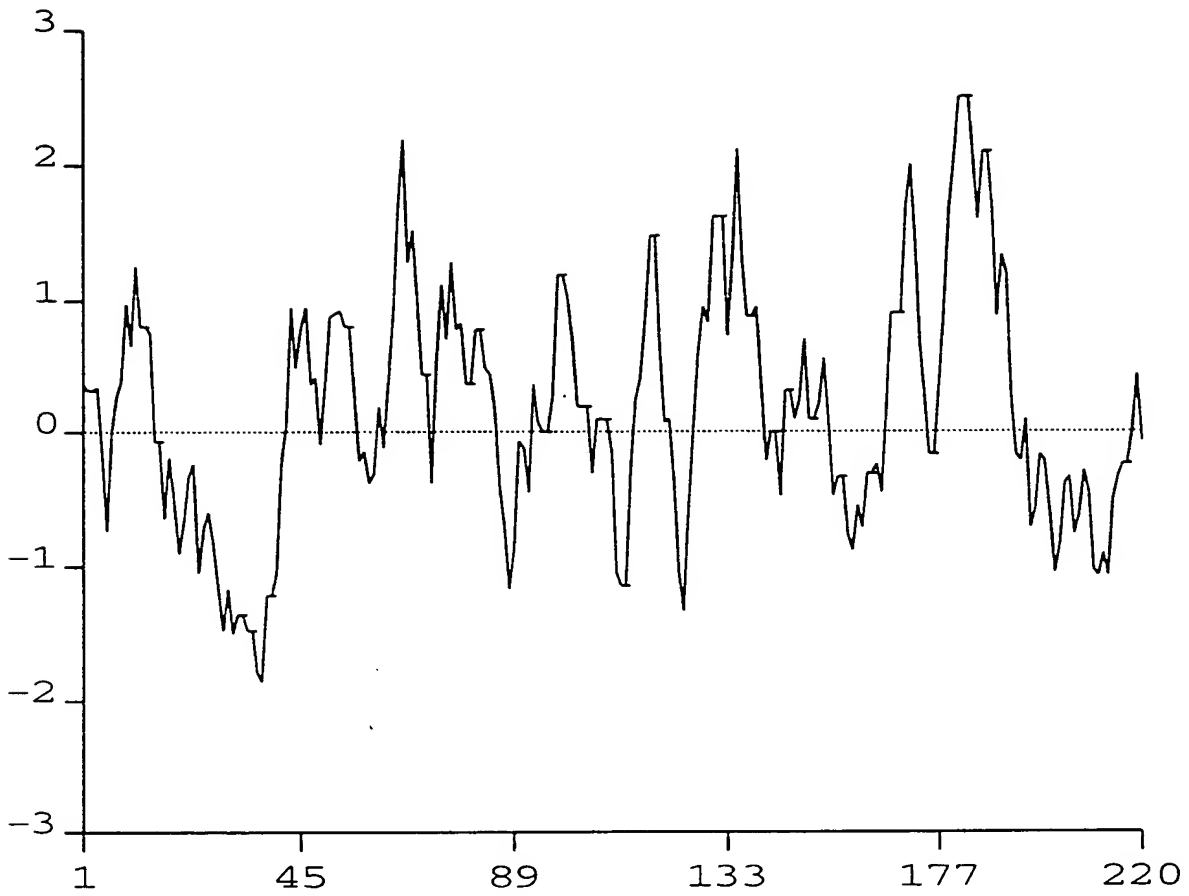


FIG. 3A

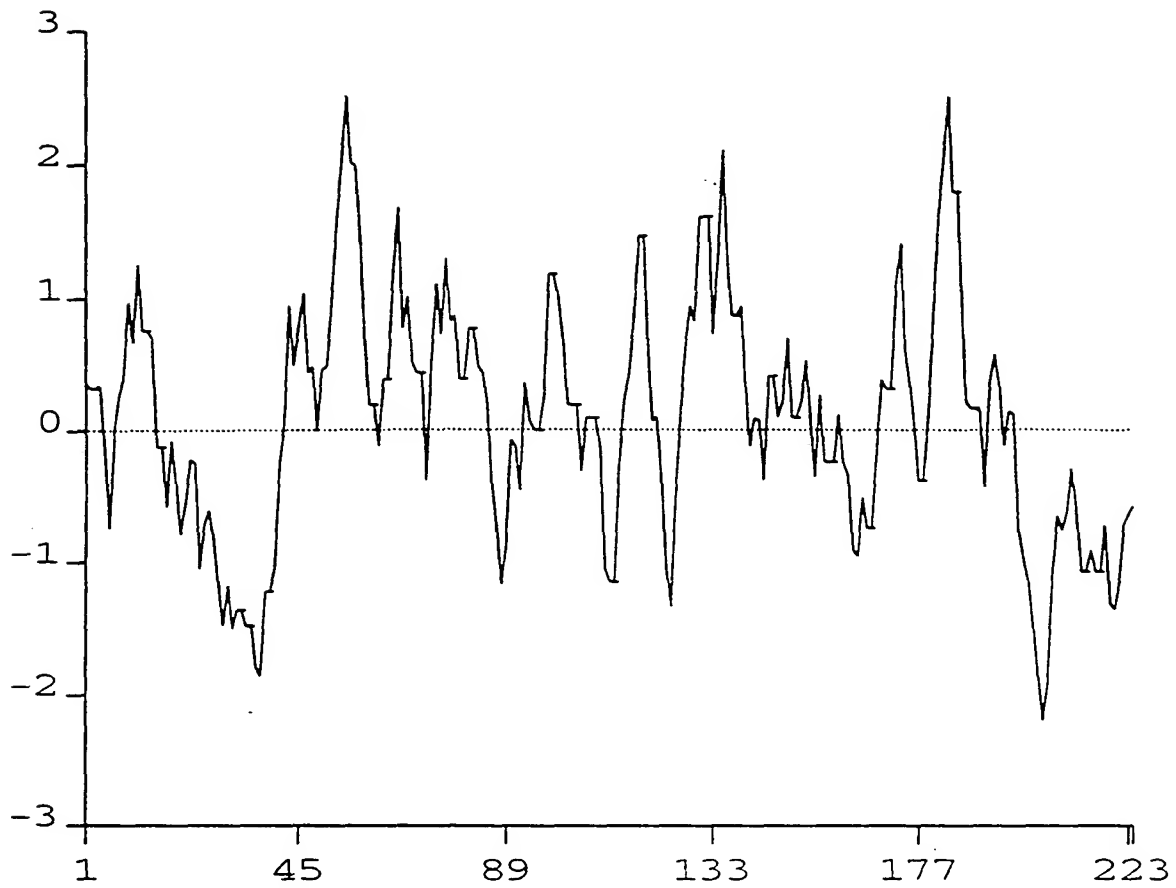


FIG. 3B